

IN THE CLAIMS

Please amend claims as shown below. Please add new claims 60-86. The following listing of the claims replaces all previous claims listings.

1. (Currently amended) A method for determining the presence of a microbial organism of interest in a sample from another organism or organisms, said method comprising:

(1) treating the sample, or a portion thereof, with at least one detectable molecular probe wherein the molecular probe or probes are peptide nucleic acid and are selected such that either:

(i) a target sequence of both the microbial organism of interest and the other organism or organisms reacts with the molecular probe in a way that produces detectable microbial organisms of interest and a detectable other organism or organisms; or

(ii) a target sequence of only the microbial organism of interest reacts with the molecular probe in a way that produces only detectable organisms of interest; ~~and~~

(2) contacting the sample, or a portion thereof, with a solid carrier to which has been immobilized an antibody such that:

if (i) applies then the antibody is chosen to be reactive only with the detectable microbial organism of interest but not reactive with the detectable other organism or organisms; but

if (ii) applies then the antibody is chosen to be generally reactive with the detectable microbial organism of interest but also may be reactive with the other organism or organisms; and

(3) determining the presence ~~or number~~ of detectable microbial organisms immobilized to the solid carrier.

2-3. (Canceled).

4. (Original) The method of claim 1, wherein the detectable molecular probe is not labeled with a detectable moiety.

5. (Previously presented) The method of claim 4, wherein the detectable molecular probe is detected using a detectable antibody that specifically binds to a complex of the detectable molecular probe and the target sequence of the microbial organism of interest.

6. (Canceled).

7. (Original) The method of claim 1, wherein the detectable molecular probe is labeled with a detectable moiety.

8. (Original) The method of claim 7, wherein the detectable moiety is selected from the group consisting of: a chromophore, a fluorochrome, a spin label, a radioisotope, an enzyme, a hapten and a chemiluminescent compound.

9-10. (Canceled).

11. (Previously presented) The method of claim 1, wherein the solid carrier is selected from the group consisting of: a particle, a bead, a microscope slide, a micro titre plate and a membrane.

12-14. (Canceled).

15. (Original) The method of claim 1, wherein the sample, or portion thereof, is treated with the detectable molecular probe or probes before being contacted with the solid carrier.

16. (Original) The method of claim 1, wherein the sample, or portion thereof, is contacted with the solid carrier before being treated with the detectable molecular probe or probes.

17. (Original) The method of claim 1, wherein the sample, or portion thereof, is simultaneously contacted with both the solid carrier and treated with the detectable molecular probe or probes.

18. (Currently amended) A method for determining presence, absence or number of a microbial organism or organisms of interest in a sample or samples, said method comprising:

(1) treating the sample or samples, or a portion thereof, with one or more detectable or independently detectable molecular probes wherein the one or more molecular probes are peptide nucleic acid and are selected such that either:

(i) the detectable probe or probes react with a target sequence of the different microbial organisms to be determined in a way that produces different detectable microbial organisms that possess the same stain; or

(ii) the independently detectable probes react with a target sequence of the different microbial organisms to be determined in a way that produces different independently detectable microbial organisms that possess an independently detectable stain; ~~and~~

(2) contacting the sample or samples, or a portion thereof, with one or more different types of coded beaded supports, wherein each different type of coded beaded support can be independently determined in a suitable particle sorter and wherein to the coded beaded supports have been immobilized one or more antibodies chosen to select a particular microbial organism or organisms such that the detectable or independently detectable organisms become selectively bound to the coded beaded supports as a result of the occurrence of specific antibody interactions;

(3) sorting the different types of coded beaded supports in a suitable particle sorter;
and

(4) determining the presence, absence, or number of detectable microbial organisms, or each of the independently detectable microbial organisms, immobilized to each different type of coded beaded support and either:

(iii) correlating the result with the particular antibody immobilized to each particle type to thereby determine the presence, absence or number of each of the different microbial organisms of interest in the sample, or portion thereof; or

(iv) correlating the result with the code for a sample source from which the sample, or portion thereof, was derived to thereby determine the presence, absence or number of each of the different organisms of interest in each different sample, or portion thereof.

19-20. (Canceled).

21. (Original) The method of claim 18, wherein the detectable molecular probe is not labeled with a detectable moiety.

22. (Currently amended) The method of claim 21, wherein the detectable molecular probe is detected using ~~a~~ detectable antibody that specifically binds to a complex of the detectable molecular probe and the target sequence of the microbial organism of interest.

23. (Canceled).

24. (Original) The method of claim 18, wherein the detectable molecular probe is labeled with a detectable moiety.

25. (Original) The method of claim 24, wherein the detectable moiety is selected from the group consisting of: a chromophore, a fluorochrome, a spin label, a radioisotope, an enzyme, a hapten and a chemiluminescent compound.

26. (Original) The method of claim 18, wherein the independently detectable probes are labeled with independently detectable fluorophores.

27-28. (Canceled).

29. (Previously presented) The method of claim 18, wherein the sample, or portion thereof, is treated with the detectable or independently detectable molecular probe or probes before being contacted with one or more different types of coded beaded supports.

30. (Previously presented) The method of claim 18, wherein the sample, or portion thereof, is contacted with one or more different types of coded beaded supports before being treated with the detectable or independently detectable molecular probe or probes.

31. (Previously presented) The method of claim 18, wherein the sample, or portion thereof, is simultaneously contacted with both one or more different types of coded beaded supports and treated with the detectable or independently detectable molecular probe or probes.

32-34. (Canceled).

35. (Currently amended) A method for determining presence, absence or number of a microbial organism or organisms in a sample, said method comprising:

(1) treating the sample, or a portion thereof, with one or more detectable or independently detectable molecular probes wherein the one or more molecular probes are peptide nucleic acid and are selected such that either:

(i) the detectable probe or probes react with a target sequence of the different microbial organisms to be determined in a way that produces different detectable microbial organisms that possess the same stain; or

(ii) the independently detectable probes react with a target sequence of the different microbial organisms to be determined in a way that produces different independently detectable microbial organisms that possess an independently detectable stain; ~~and~~

(2) contacting the sample, or a portion thereof, with a solid carrier array to which antibodies have been immobilized at unique identifiable locations such that the different detectable microbial organisms or the different independently detectable microbial organisms are selectively bound to the locations on the array as a result of the occurrence of specific antibody interactions; ~~and~~

(3) determining the presence, absence or number of the detectable or independently detectable microbial organisms immobilized at the many different locations of the array; and

(4) correlating the result with the particular antibody immobilized to each location on the array to thereby determine the presence, absence or number of the different microbial organisms of interest in the sample.

36-37. (Canceled).

38. (Original) The method of claim 35, wherein the detectable molecular probe is not labeled with a detectable moiety.

39. (Previously presented) The method of claim 38, wherein the detectable molecular probe is detected using a detectable antibody that specifically binds to a complex of the detectable molecular probe and the target sequence of the microbial organism of interest.

40. (Canceled).

41. (Original) The method of claim 35, wherein the detectable molecular probe is labeled with a detectable moiety.

42. (Original) The method of claim 41, wherein the detectable moiety is selected from the group consisting of: a chromophore, a fluorochrome, a spin label, a radioisotope, an enzyme, a hapten and a chemiluminescent compound.

43. (Original) The method of claim 35, wherein the independently detectable probes are labeled with independently detectable fluorophores.

44-45. (Canceled).

46. (Original) The method of claim 35, wherein the sample is treated with the detectable or independently detectable molecular probe or probes before being contacted with the solid carrier.

47. (Original) The method of claim 35, wherein the sample is contacted with the solid carrier before being treated with the detectable or independently detectable molecular probe or probes.

48. (Original) The method of claim 35, wherein the sample is simultaneously contacted with both the solid carrier and treating with the detectable or independently detectable molecular probe or probes.

49-59. (Canceled).

60. (New) A method for determining the presence of a microbial organism of interest in a sample from another organism or organisms, said method comprising:

(1) treating the sample, or a portion thereof, with at least one detectable molecular probe wherein the molecular probe or probes are peptide nucleic acid and are selected such that either:

(i) a target sequence of both the microbial organism of interest and the other organism or organisms reacts with the molecular probe in a way that produces detectable microbial organisms of interest and a detectable other organism or organisms;
or

(ii) a target sequence of only the microbial organism of interest reacts with the molecular probe in a way that produces only detectable organisms of interest;

(2) contacting the sample, or a portion thereof, with a solid carrier to which has been immobilized an antibody such that:

if (i) applies then the antibody is chosen to be reactive only with the detectable microbial organism of interest but not reactive with the detectable other organism or organisms; but

if (ii) applies then the antibody is chosen to be generally reactive with the detectable microbial organism of interest but also may be reactive with the other organism or organisms; and

(3) determining the presence of detectable microbial organisms immobilized to the solid carrier,

wherein the microbial organism of interest is consisting of a cell, a bacterium, a virus, a yeast, a fungus, another unicellular organism and a multicellular organism.

61. (New) The method of claim 60, wherein the detectable molecular probe is not labeled with a detectable moiety.

62. (New) The method of claim 61, wherein the detectable molecular probe is detected using a detectable antibody that specifically binds to a complex of the detectable molecular probe and the target sequence of the microbial organism of interest.

63. (New) The method of claim 60, wherein the detectable molecular probe is labeled with a detectable moiety.

64. (New) The method of claim 63, wherein the detectable moiety is selected from the group consisting of: a chromophore, a fluorochrome, a spin label, a radioisotope, an enzyme, a hapten and a chemiluminescent compound.

65. (New) The method of claim 60, wherein the solid carrier is selected from the group consisting of: a particle, a bead, a microscope slide, a micro titre plate and a membrane.

66. (New) The method of claim 60, wherein the sample, or portion thereof, is treated with the detectable molecular probe or probes before being contacted with the solid carrier.

67. (New) The method of claim 60, wherein the sample, or portion thereof, is contacted with the solid carrier before being treated with the detectable molecular probe or probes.

68. (New) The method of claim 60, wherein the sample, or portion thereof, is simultaneously contacted with both the solid carrier and treated with the detectable molecular probe or probes.

69. (New) A method for determining presence, absence or number of a microbial organism or organisms of interest in a sample or samples, said method comprising:

(1) treating the sample or samples, or a portion thereof, with one or more detectable or independently detectable molecular probes wherein the one or more molecular probes are peptide nucleic acid and are selected such that either:

(i) the detectable probe or probes react with a target sequence of the different microbial organisms to be determined in a way that produces different detectable microbial organisms that possess the same stain; or

(ii) the independently detectable probes react with a target sequence of the different microbial organisms to be determined in a way that produces different independently detectable microbial organisms that possess an independently detectable stain;

(2) contacting the sample or samples, or a portion thereof, with one or more different types of coded beaded supports, wherein each different type of coded beaded support can be independently determined in a suitable particle sorter and wherein to the coded beaded supports have been immobilized one or more antibodies chosen to select a particular microbial organism or organisms such that the detectable or independently detectable organisms become selectively bound to the coded beaded supports as a result of the occurrence of specific antibody interactions;

(3) sorting the different types of coded beaded supports in a suitable particle sorter;
and

(4) determining the presence, absence, or number of detectable microbial organisms, or each of the independently detectable microbial organisms, immobilized to each different type of coded beaded support and either:

(iii) correlating the result with the particular antibody immobilized to each particle type to thereby determine the presence, absence or number of each of the different microbial organisms of interest in the sample, or portion thereof; or

(iv) correlating the result with the code for a sample source from which the sample, or portion thereof, was derived to thereby determine the presence, absence or number of each of the different organisms of interest in each different sample, or portion thereof,

wherein the microbial organism of interest is selected from the group consisting of a cell, a bacterium, a virus, a yeast, a fungus, another unicellular organism and a multicellular organism.

70. (New) The method of claim 69, wherein the detectable molecular probe is not labeled with a detectable moiety.

71. (New) The method of claim 70, wherein the detectable molecular probe is detected using a detectable antibody that specifically binds to a complex of the detectable molecular probe and the target sequence of the microbial organism of interest.

72. (New) The method of claim 69, wherein the detectable molecular probe is labeled with a detectable moiety.

73. (New) The method of claim 72, wherein the detectable moiety is selected from the group consisting of: a chromophore, a fluorochrome, a spin label, a radioisotope, an enzyme, a hapten and a chemiluminescent compound.

74. (New) The method of claim 69, wherein the independently detectable probes are labeled with independently detectable fluorophores.

75. (New) The method of claim 69, wherein the sample, or portion thereof, is treated with the detectable or independently detectable molecular probe or probes before being contacted with one or more different types of coded beaded supports.

76. (New) The method of claim 69, wherein the sample, or portion thereof, is contacted with one or more different types of coded beaded supports before being treated with the detectable or independently detectable molecular probe or probes.

77. (New) The method of claim 69, wherein the sample, or portion thereof, is simultaneously contacted with both one or more different types of coded beaded supports and treated with the detectable or independently detectable molecular probe or probes.

78. (New) A method for determining presence, absence or number of a microbial organism or organisms in a sample, said method comprising:

(1) treating the sample, or a portion thereof, with one or more detectable or independently detectable molecular probes wherein the one or more molecular probes are peptide nucleic acid and are selected such that either:

(i) the detectable probe or probes react with a target sequence of the different microbial organisms to be determined in a way that produces different detectable microbial organisms that possess the same stain; or

(ii) the independently detectable probes react with a target sequence of the different microbial organisms to be determined in a way that produces different independently detectable microbial organisms that possess an independently detectable stain;

(2) contacting the sample, or a portion thereof, with a solid carrier array to which antibodies have been immobilized at unique identifiable locations such that the different detectable microbial organisms or the different independently detectable microbial organisms are selectively bound to the locations on the array as a result of the occurrence of specific antibody interactions;

(3) determining the presence, absence or number of the detectable or independently detectable microbial organisms immobilized at the many different locations of the array;
and

(4) correlating the result with the particular antibody immobilized to each location on the array to thereby determine the presence, absence or number of the different microbial organisms of interest in the sample,

wherein the microbial organism of interest is selected from the group consisting of a cell, a bacterium, a virus, a yeast, a fungus, another unicellular organism and a multicellular organism.

79. (New) The method of claim 78, wherein the detectable molecular probe is not labeled with a detectable moiety.

80. (New) The method of claim 79, wherein the detectable molecular probe is detected using a detectable antibody that specifically binds to a complex of the detectable molecular probe and the target sequence of the microbial organism of interest.

81. (New) The method of claim 78, wherein the detectable molecular probe is labeled with a detectable moiety.

82. (New) The method of claim 81, wherein the detectable moiety is selected from the group consisting of: a chromophore, a fluorochrome, a spin label, a radioisotope, an enzyme, a hapten and a chemiluminescent compound.

83. (New) The method of claim 78, wherein the independently detectable probes are labeled with independently detectable fluorophores.

84. (New) The method of claim 78, wherein the sample is treated with the detectable or independently detectable molecular probe or probes before being contacted with the solid carrier.

85. (New) The method of claim 78, wherein the sample is contacted with the solid carrier before being treated with the detectable or independently detectable molecular probe or probes.

In Re Application Of:

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Application No.: 09/966,658

Filed: November 29, 2001

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Attorney Docket No.: BP-0002-1 US

86. (New) The method of claim 78, wherein the sample is simultaneously contacted with both the solid carrier and treating with the detectable or independently detectable molecular probe or probes.